

Fig. 1.

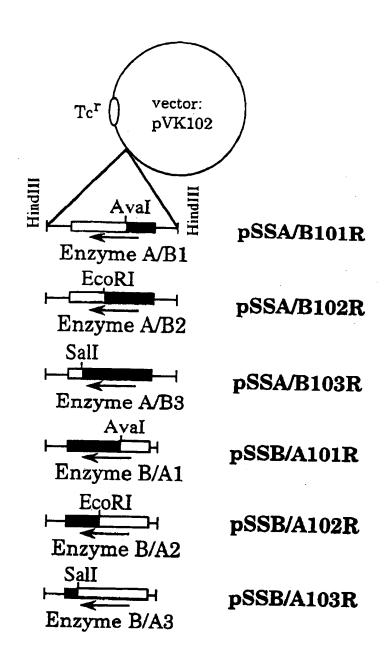


Fig. 2.

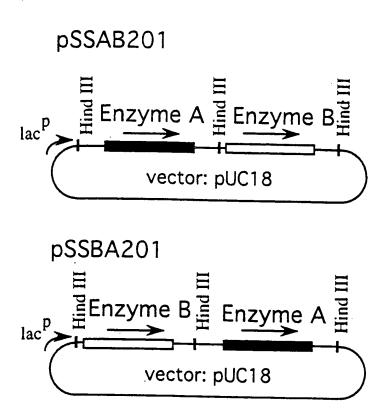
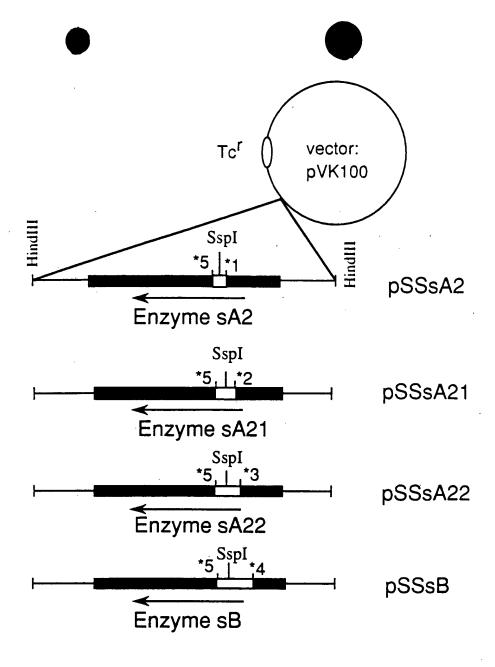


Fig. 3.



Recombination site

*1 : amino acid residue No. 135 of mature Enzyme A

*2 : amino acid residue No. 128 of mature Enzyme A

*3: amino acid residue No. 125 of mature Enzyme A

*4 : amino acid residue No. 95 of mature Enzyme A

*5: amino acid residue No. 180 of mature Enzyme B, which nucleotide sequence of Aval site encodes

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Fig. 4.

mn 3 = 0 0

Enzyme A 1 : QVTPVTDELL ANPPAGEWIS YGQNQENYRH SPLTQITTEN VGQLQLVWAR GMQPGKVQVT Enzyme B 1 : QVTPITDELL ANPPAGEWIN YGRNQENYRH SPLTQITADN VGQLQLVWAR GMEAGAVQVT 61 : PLIHDGVMYL ANPGDVIQAI DAKTGDLIWE HRRQLPNIAT LNSFGEPTRG MALYGTNVYF 61 : PMIHDGVMYL ANPGDVIQAL DAQTGDLIWE HRRQLPAVAT LNAQGDRKRG VALYGTSLYF 121 : VSWDNHLVAL DTATGQVTFD VDRGQGED-M VSNSSGPIVA NGVIVAGSTC QYSPFGCFVS Aval ***** ** * **** ** * ** ** **** ******* *** * SSWDNHLIAL DMETGQVVFD VERGSGEDGL TSNTTGPIVA NGVIVAGSTC QYSPYGCFIS 180 : GHDSATGEEL WRNYFIPRAG EEGDETWGND YEARWMTGAW GQITYDPVTN LVHYGSTAVG ******** *** *** 181 : GHDSATGEEL WRNHFIPQPG EEGDETWGND FEARWMTGVW GQITYDPVTN LVFYGSTGVG 240 : PASETQRGTP GGTLYGTNTR FAVRPDTGEI VWRHQTLPRD NWDQECTFEM MVTNVDVQPS PASETQRGTP GGTLYGTNTR FAVRPDTGEI VWRHQTLPRD NWDQECTFEM MVANVDVQPS EcoRI 300 : TEMEGLQSIN PNAATGERRV LTGVPCKTGT MWQFDAETGE FLWARDTNYQ NMIESIDENG 301 : AEMEGLRAIN PNAATGERRV LTGAPCKTGT MWSFDAASGE FLWARDTNYT NMIASIDETG 360 : IVTVNEDAIL KELDVEYDVC PTFLGGRDWP SAALNPDSGI YFIPLNNVCY DMMAVDQEFT ******* * ******** ****** ****** ** ** *** ** * * **** 361 : LVTVNEDAVL KELDVEYDVC PTFLGGRDWS SAALNPDTGI YFLPLNNACY DIMAVDQEFS SalI 420 : SMDVYNTSNV TKLPPGKDMI GRIDAIDIST GRTLWSVERA AANYSPVLST GGGVLFNGGT 421 : ALDVYNTSAT AKLAPGFENM GRIDAIDIST GRTLWSAERP AANYSPVLST AGGVVFNGGT 480 : DRYFRALSQE TGETLWQTRL ATVASGQAIS YEVDGMQYVA IAGGGVSYGS GLNSALAGER ********* 481 : DRYFRALSQE TGETLWQARL ATVATGQAIS YELDGVQYIA IGAGGLTYGT QLNAPLA-EA 540 : VDSTAIGNAV YVFALPQ 540 : IDSTSVGNAI YVFALPO

Fig. 5.

^{* :} Nucleotide sequences encoding these regions are the restriction sites for Aval, EcoRI, and Sall which were used for constructing chimera genes shown in Fig. 2.

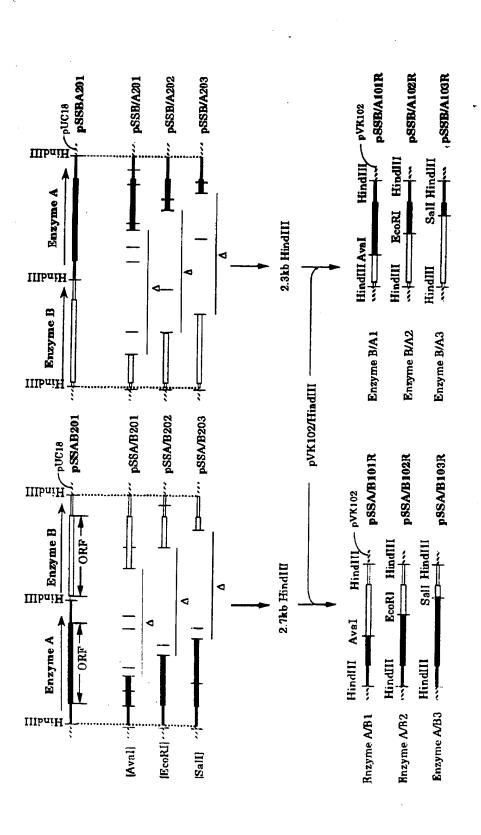
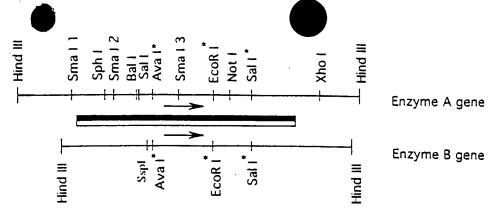


Fig. 6.



*: Aval, EcoRI, Sall sites used for constructing chimera genes shown in Figs. 2 and 6.

Fig. 7.

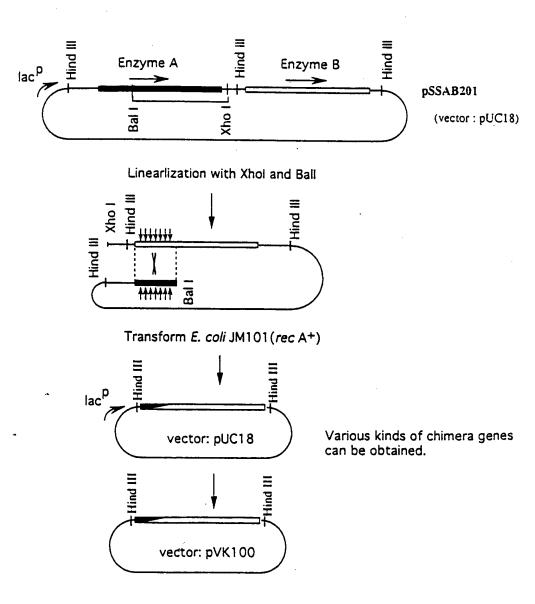


Fig. 8.

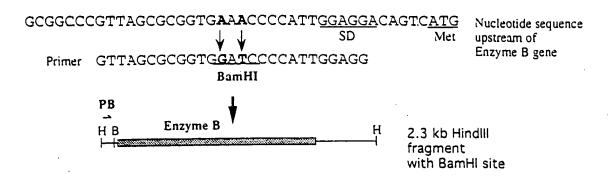


Fig. 9.

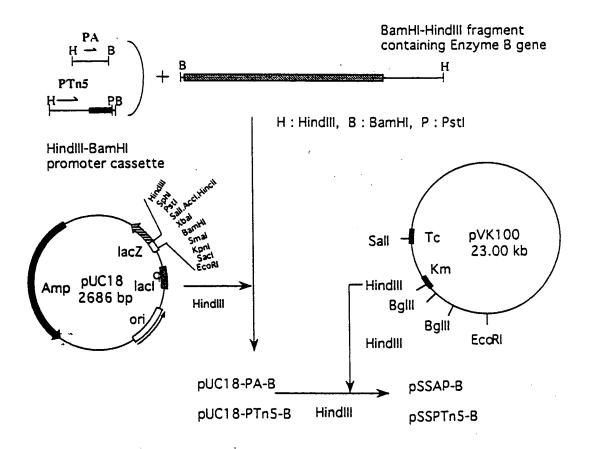
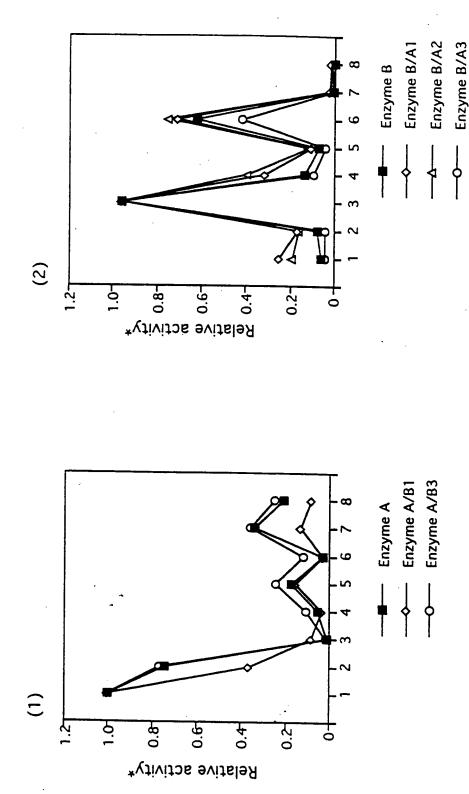


Fig. 10.

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1. n-Propanol, 2. Isopropanol, 3. D-Glucose, 4. L-Sorbosone 5. D-Sorbitol, 6. D-Mannitol, 7. L-Sorbose, 8. D-Fructose

*Enzyme activity was normalized relative to activity for n-propanol (1), or D-glucose (2). Enzyme A/B2 was excepted because of its low expression in P. putida.

Fig. 11.